



1600

RAW SEQUENCE LISTINGPATENT APPLICATION: **US/09/937,555B**DATE: 09/10/2003

TIME: 08:48:35

Input Set : A:\Hookworm.ST25.txt

Output Set: N:\CRF4\09102003\I937555B.raw

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3 <110> APPLICANT: Cappello, Michael
         Chadderdon, Robert
         Del Valle, Antonio
        Harrison, Lisa
 8 <120> TITLE OF INVENTION: Hookworm Platelet Inhibitor
10 <130> FILE REFERENCE: OCR-1001
12 <140> CURRENT APPLICATION NUMBER: 09/937,555B
13 <141> CURRENT FILING DATE: 2000-03-30
15 <150> PRIOR APPLICATION NUMBER: US 60/127,239
16 <151> PRIOR FILING DATE: 1999-03-30
18 <160> NUMBER OF SEQ ID NOS: 6
20 <170> SOFTWARE: PatentIn version 3.2
22 <210> SEQ ID NO: 1
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23 <211> LENGTH: 631
24 <212> TYPE: DNA
25 <213> ORGANISM: Ancylostoma caninum
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32 atgagaacta tggtttacga ctgtaccttg gaagaaaagg cctacaaatc ggctgaaaaa
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34 tgcagtgagg aaccetegag tgaggaagaa aacgtagatg ttttcagege agetactett
36 aatatteegt tagaggeegg taatteatgg tggagegaga ttttegaact gegaggaaag
38 gtctacaaca aaaatggcaa aacatcgaac attgctaata tggtttggga cagtcatgat
40 aagcttggtt gegeagttgt tgactgetee ggaaaaacae atgtagtetg ecaataeggg
42 ccagaagcaa aaggtgatgg gaagacaatt tacgaagagg gtgcaccatg ctcacgatgc
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44 agtgattacg gggcaggtgt cacctgcgac gacgactggc agaatttgct ctgcattggt
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59 <222> LOCATION: (1)..(40)
60 <223> OTHER INFORMATION: residues 1-40 native; remaining residues deduced amino acid
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63 <400> SEQUENCE: 2
65 Glu Gly Asp Tyr Ser Leu Cys Gln Gln Arg Glu Lys Leu Asp Asp
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69 Met Arg Glu Met Phe Thr Glu Leu His Asn Gly Tyr Arg Ala Ala Phe
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73 Ala Arg Asn Tyr Lys Thr Ser Lys Met Arg Thr Met Val Tyr Asp Cys
```

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RAW SEQUENCE LISTING PATENT APPLICATION: US/09/937,555B Input Set : A:\Hookworm.ST25.txt Output Set: N:\CRF4\09102003\I937555B.raw 77 Thr Leu Glu Glu Lys Ala Tyr Lys Ser Ala Glu Lys Cys Ser Glu Glu 81 Pro Ser Ser Glu Glu Glu Asn Val Asp Val Phe Ser Ala Ala Thr Leu 78 50 85 Asn Ile Pro Leu Glu Ala Gly Asn Ser Trp Trp Ser Glu Ile Phe Glu 89 Leu Arg Gly Lys Val Tyr Asn Lys Asn Gly Lys Thr Ser Asn Ile Ala 93 Asn Met Val Trp Asp Ser His Asp Lys Leu Gly Cys Ala Val Val Asp 90 100 97 Cys Ser Gly Lys Thr His Val Val Cys Gln Tyr Gly Pro Glu Ala Lys 101 Gly Asp Gly Lys Thr Ile Tyr Glu Glu Gly Ala Pro Cys Ser Arg Cys 105 Ser Asp Tyr Gly Ala Gly Val Thr Cys Asp Asp Trp Gln Asn Leu 106 109 Leu Cys Ile Gly 180 110 113 <210> SEQ ID NO: 3 114 <211> LENGTH: 6 115 <212> TYPE: PRT 116 <213> ORGANISM: Ancylostoma caninum 118 <400> SEQUENCE: 3 120 Glu Gly Asp Tyr Ser Leu 121 1 121 1 124 <210> SEQ ID NO: 4 125 <211> LENGTH: 6 126 <212> TYPE: PRT 127 <213> ORGANISM: Ancylostoma caninum 129 <400> SEQUENCE: 4 131 Glu Gly Asp Tyr Ser Leu 132 1 135 <210> SEQ ID NO: 5 136 <211> LENGTH: 8 137 <212> TYPE: PRT 138 <213> ORGANISM: Ancylostoma caninum 140 <400> SEQUENCE: 5 142 Thr Ser Asn Ile Ala Asn Met Val . 5 143 1 146 <210> SEQ ID NO: 6 147 <211> LENGTH: 8 148 <212> TYPE: PRT 149 <213> ORGANISM: Ancylostoma caninum 151 <400> SEQUENCE: 6 153 Thr Ser Asn Ile Ala Asn Met Val 154 1

VERIFICATION SUMMARY

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